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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/050,054

DATE: 02/11/2002
TIME: 10:17:49

Input Set : A:\474.app
Output Set: N:\CRF3\02112002\J050054.raw

4 <110> APPLICANT: Alnemri, Emad S.
6 <120> TITLE OF INVENTION: INHIBITION OF INTERLEUKIN-1-BETA
7 SECRETION BY CARD PROTEINS
9 <130> FILE REFERENCE: 480140.474
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/050,054
12 <141> CURRENT FILING DATE: 2002-01-16
14 <160> NUMBER OF SEQ ID NOS: 8
16 <170> SOFTWARE: FastSEQ for Windows Version 4.0
18 <210> SEQ ID NO: 1
19 <211> LENGTH: 97
20 <212> TYPE: PRT
21 <213> ORGANISM: Homo sapiens
23 <400> SEQUENCE: 1
24 Met Ala Asp Lys Val Leu Lys Glu Lys Arg Lys Leu Phe Ile His Ser
25 1 5 10 15
26 Met Gly Glu Gly Thr Ile Asn Gly Leu Leu Asp Glu Leu Leu Gln Thr
27 20 25 30
28 Arg Val Leu Asn Gln Glu Glu Met Glu Lys Val Lys Arg Glu Asn Ala
29 35 40 45
30 Thr Val Met Asp Lys Thr Arg Ala Leu Ile Asp Ser Val Ile Pro Lys
31 50 55 60
32 Gly Ala Gln Ala Cys Gln Ile Cys Ile Thr Tyr Ile Cys Glu Glu Asp
33 65 70 75 80
34 Ser Tyr Leu Ala Glu Thr Leu Gly Leu Ser Ala Gly Pro Ile Pro Gly
35 85 90 95
36 Asn
40 <210> SEQ ID NO: 2
41 <211> LENGTH: 294
42 <212> TYPE: DNA
43 <213> ORGANISM: Homo sapiens
45 <400> SEQUENCE: 2
46 atggccgaca aggtcctgaa ggagaagaga aagctgttta tccattccat gggtaaggt 60
47 acaataaaatg gcttactgga tgaattatta cagacaaggg tgctgaacca ggaagagatg 120
48 gagaaagtaa aacgtaaaaa tgctacagtt atggataaga cccgagctt gattgactcc 180
49 gtattccgaa aagggcaca ggcattccaa atttgcattca catacatttg tgaagaagac 240
50 agttacctgg cagagacgt gggactctca gcaggccga tacctggaaa ttag 294
52 <210> SEQ ID NO: 3
53 <211> LENGTH: 90
54 <212> TYPE: PRT
55 <213> ORGANISM: Homo sapiens
57 <400> SEQUENCE: 3
58 Met Ala Asp Gln Leu Leu Arg Lys Lys Arg Arg Ile Phe Ile His Ser
59 1 5 10 15

ENTITLED

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60 Val Gly Ala Gly Thr Ile Asn Ala Leu Leu Asp Cys Leu Leu Glu Asp
61 20 25 30
62 Glu Val Ile Ser Gln Glu Asp Met Asn Lys Val Arg Asp Glu Asn Asp
63 35 40 45
64 Thr Val Met Asp Lys Ala Arg Val Leu Ile Asp Leu Val Thr Gly Lys
65 50 55 60
66 Gly Pro Lys Ser Cys Cys Lys Phe Ile Lys His Leu Cys Glu Glu Asp
67 65 70 75 80
68 Pro Gln Leu Ala Ser Lys Met Gly Leu His
69 85 90
72 <210> SEQ ID NO: 4
73 <211> LENGTH: 273
74 <212> TYPE: DNA
75 <213> ORGANISM: Homo sapiens
77 <400> SEQUENCE: 4
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79 acaataaatg ccttgcgttga ttgccttattt gaggatgaag ttatttagccca ggaagacatg 120
80 aacaaagtga gagatgaaaa tgacactgtc atggataagg ctcgagtctt gattgacctt 180
81 gttactggaa aaggacccaa gtcttgcgtc aaatttatca agcatctctg tgaagaagac 240
82 cctcaacttg cctcaaaatg gggtttgcac taa 273
84 <210> SEQ ID NO: 5
85 <211> LENGTH: 103
86 <212> TYPE: PRT
87 <213> ORGANISM: Homo sapiens
89 <400> SEQUENCE: 5
90 Met Ala Asp Lys Val Leu Lys Glu Lys Arg Lys Leu Phe Ile Arg Ser
91 1 5 10 15
92 Met Gly Glu Gly Thr Ile Asn Gly Leu Leu Asp Glu Leu Gln Thr
93 20 25 30
94 Arg Val Leu Asn Lys Glu Glu Met Glu Lys Val Lys Arg Glu Asn Ala
95 35 40 45
96 Thr Val Met Asp Lys Thr Arg Ala Leu Ile Asp Ser Val Ile Pro Lys
97 50 55 60
98 Gly Ala Gln Ala Cys Gln Ile Cys Ile Thr Tyr Ile Cys Glu Glu Asp
99 65 70 75 80
100 Ser Tyr Leu Ala Gly Thr Leu Gly Leu Ser Ala Asp Gln Thr Ser Gly
101 85 90 95
102 Asn Tyr Leu Asn Met Gln Asp
103 100
106 <210> SEQ ID NO: 6
107 <211> LENGTH: 429
108 <212> TYPE: DNA
109 <213> ORGANISM: Homo sapiens
111 <400> SEQUENCE: 6
112 atggccgaca aggttctgaa ggagaagaga aagctttta tccgttccat gggtaagg 60
113 acaataaatg gcttactgttga tgaatttta cagacaagg tgctgaacaa ggaagagatg 120
114 gagaaagtta aacgtgaaaa tgctacagt atggataaga cccgagctt gattgactcc 180
115 gttattccga aaggggcaca ggcatgcacat tttgcacatacattt tgaagaagac 240
116 agttacctgg caggacgct gggactctca gcagatcaaa catctggaaa ttaccttaat 300

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117 atgcaagact ctcaaggagt actttcttcc tttccagctc ctcaggcagt gcaggacaac 360
118 ccagctatgc ccacatcctc aggctcagaa gggaatgtca agcttgctc cctagaagaa 420
119 gctcaaagg 429
121 <210> SEQ ID NO: 7
122 <211> LENGTH: 430
123 <212> TYPE: DNA
124 <213> ORGANISM: Homo sapiens
126 <400> SEQUENCE: 7
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128 acaataaaatg gcttaactgga tgaattatta cagacaaggg tgctgaacca ggaagagatg 120
129 gagaaagtaa aacgtgaaaa tgctacagtt atggataaga cccgagctt gattgactcc 180
130 gttattccga aaggggcaca ggcattgcca atttgcatca catacatgg tgaagaagac 240
131 agttacctgg cagagacgct gggactctca gcaggtccga tacctggaaa ttagcttagt 300
132 acacaagact cccaaattact attttcttcc ttcccagctc ttcaggcagt gcgaggacaa 360
133 cccagctatg cccacatgct caagcccaga aggagaatc aagcttgct ttctagaaga 420
134 cgctcaagg 430
136 <210> SEQ ID NO: 8
137 <211> LENGTH: 52
138 <212> TYPE: DNA
139 <213> ORGANISM: Homo sapiens
141 <400> SEQUENCE: 8
142 atggccgggt agtaaacaga caaggcagca ggtaaggta taggtccgat ag 52

VERIFICATION SUMMARY

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L:11 M:270 C: Current Application Number differs, Wrong Format